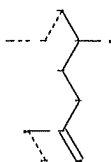


chain nodes :
 1 2 3 4 5 6 9 10 11 13 15
 Chain bonds :
 1-2 1-4 2-3 4-5 4-15 5-6 6-9 9-10 9-11 11-13
 exact/norm bonds :
 1-2 1-4 2-3 4-15 6-9 9-10 11-13
 exact bonds :
 4-5 5-6 9-11



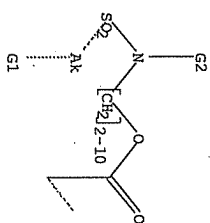
G1:CF2,CF3

G2:H,Ak

Match level :
 1:CLASS 2:CLASS 3:CLASS 4:CLASS 5:CLASS 6:CLASS 9:CLASS 10:CLASS 11:CLASS
 13:CLASS 15:CLASS

L1 STRUCTURE UPLOADED

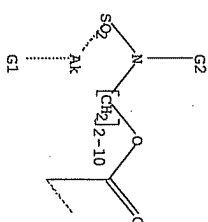
=> d l1
 l1 HAS NO ANSWERS
 l1 STR



G1 CF2,CF3
 G2 H,Ak

Structure attributes must be viewed using STN Express query preparation.

=> d que stat l1
 l1 STR



G1 CF2,CF3
 G2 H,Ak

Structure attributes must be viewed using STN Express query preparation.

=> s file=hcapius abb-on l1
 FILE 'HCAPLUS' IS NOT VALID IN THE CURRENT FILE ENVIRONMENT
 REGISTRY
 ENTER A FILE NAME OR (ALL):hcapius
 FILE 'HCAPLUS' IS NOT VALID IN THE CURRENT FILE ENVIRONMENT
 REGISTRY
 ENTER A FILE NAME OR (ALL):
 ENTER A FILE NAME OR (ALL):file hcapius
 FILE 'FILE HCAPLUS' IS NOT VALID IN THE CURRENT FILE ENVIRONMENT
 REGISTRY
 ENTER A FILE NAME OR (ALL):all
 SAMPLE SEARCH INITIATED 20:38:55 FILE 'REGISTRY'
 SAMPLE SCREEN SEARCH COMPLETED - 233 TO ITERATE

100.0% PROCESSED 233 ITERATIONS
 INCOMPLETE SEARCH (SYSTEM LIMIT EXCEEDED)
 SEARCH TIME: 00.00.01 50 ANSWERS

FULL FILE PROJECTIONS: ONLINE **COMPLETE**
 BATCH 3745 TO 5575
 PROJECTED ITERATIONS: 1064 TO 2136
 PROJECTED ANSWERS:
 l2 50 SEA SSS SAM l1

=> s l1
 REGISTRY INITIATED
 Substance data SEARCH and crossover from CAS REGISTRY in progress...
 Use DISPLAY HITSTR (or HITSTR) to directly view retrieved structures.

SAMPLE SEARCH INITIATED 20:46:25 FILE 'REGISTRY'
 SAMPLE SCREEN SEARCH COMPLETED - 233 TO ITERATE
 100.0% PROCESSED 233 ITERATIONS 50 ANSWERS